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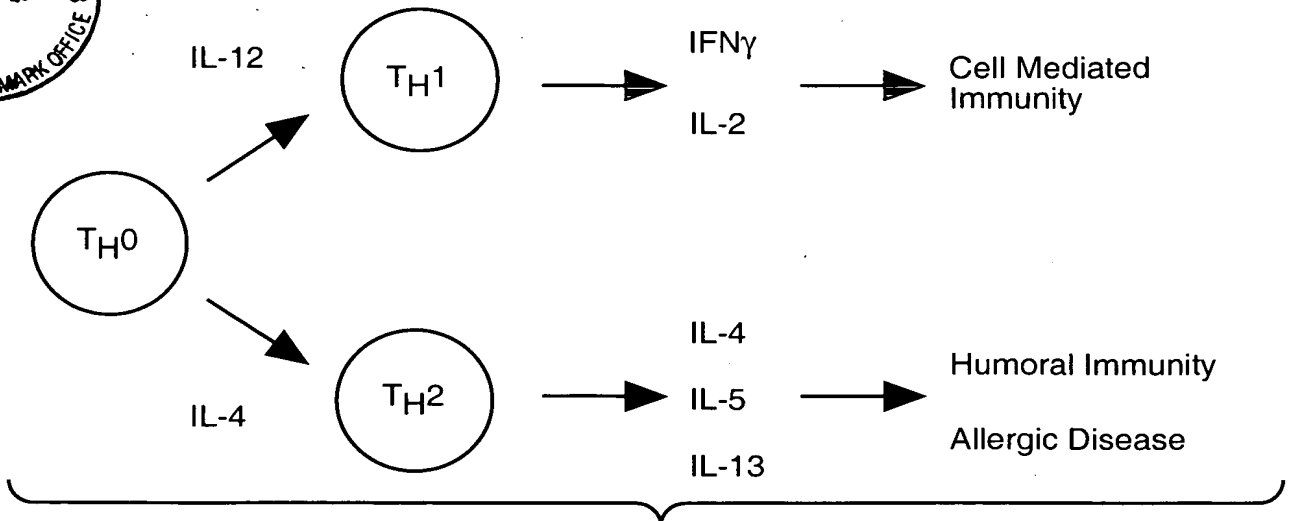


FIG. 1

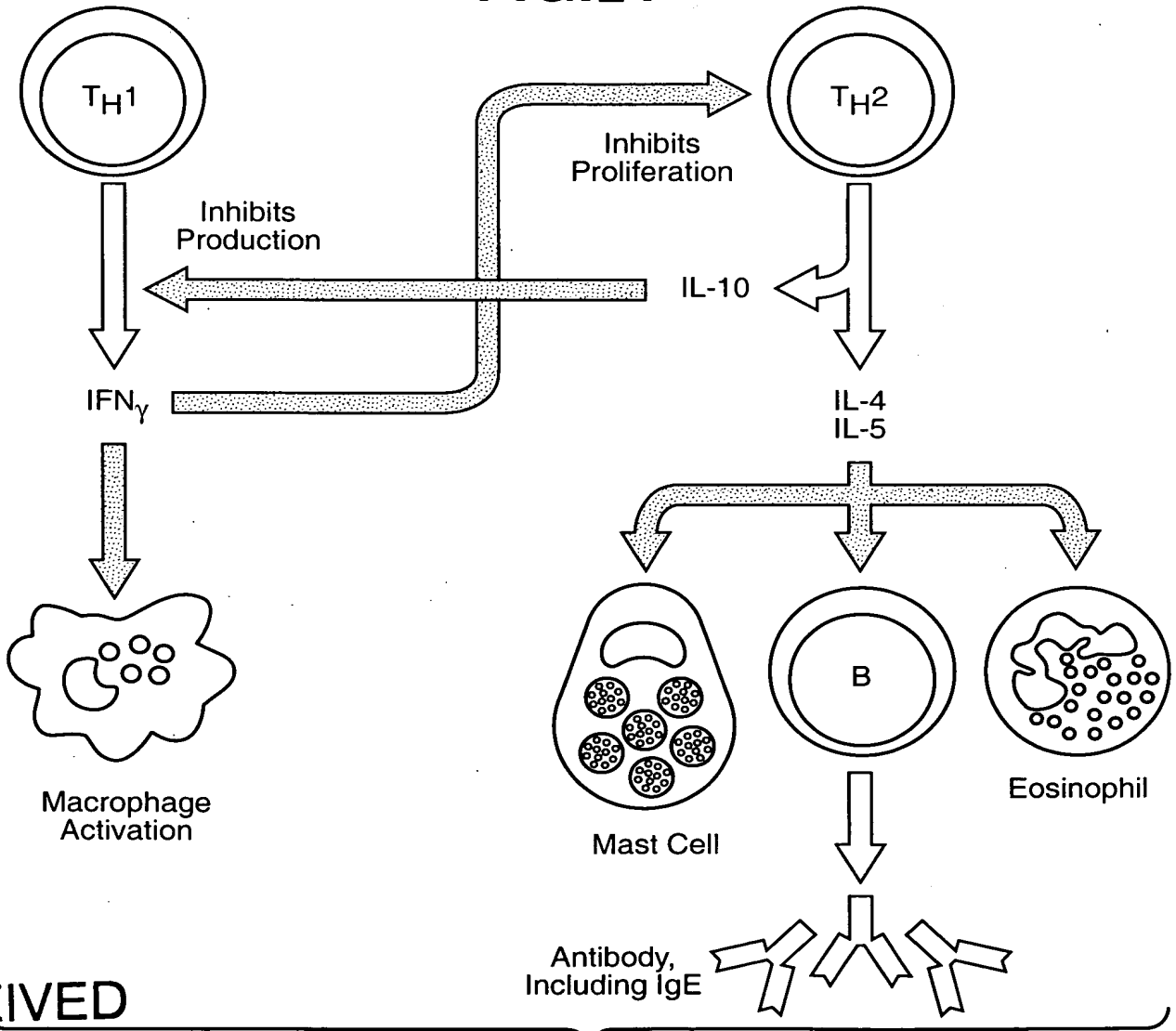


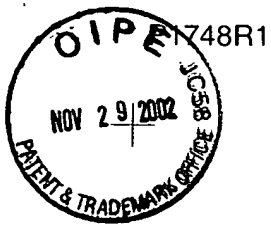
FIG. 2

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MRGGRGAPFWLWPLPKLALLPLLWVLFQRTRPQGSAGPLQCYGVGPLGDLNCSWEPLGD
LGAPSELHLQSQKYRSNKTQTVAVAAGRSWVAIPREQLTMSDKLLVWGTKAGQPLWPPV
FVNLETQMKPNAPRLGPDVDFSEDDPLEATVHWAPPTWPSHKVLICQFHYRRCQEAAWT
LLEPELKTIPLTPEIQDLELATGYKVYGRCRMEKEEDLWGEWSPILSFQTPPSAPKDV
WVSGNLCGTPGGEEPLLLWKAPGPCVQVSYKVWFVVGRELSPEGITCCCSLIPSGAEW
ARVSAVNATSWEPLTNLSLVCLDSASAPRSVAVSSIAGSTELLVTWQPGPGEPLHVVD
WARDGDPLEKLNWVRLPPGNLSALLPGNFTVGVPYRITVTAVSASGLASASSVWGFREE
LAPLVGPTLWRLQDAPPGTPAIAWGEVPRHQLRGHLTHYTLCAQSGTSPSVCNMVSGNT
QSVTLPLDLPWGPCELWVTASTIAGQGGPILRLHLPDNTLRWKVLPGLFLWGLFLLG
CGLSLATSGRCYHLRHKVLPRVWWEKVPDPANSSSGQPHMEQVPEAQPLGDLPILEVEE
MEPPPVMESSQPAQATAPLDSGYEKHFLPTPEELGLLGPPRPQVLA

APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

FIG._3

MNRLRVARLTPLELLLSLMSLLLGTTRPHGSPGLQCYSVGPLGILNCSWEPLGDLETTPV
LYHQSQKYHPNRVWEVKVPSKQSWVTIPREQFTMADKLLIWGTQKGRPLWSSSVNLETQ
MKPDTPQIFSQVDISEEATLEATVQWAPPVWPPQKALTCQFRYKECQAEAWTRLEPQLKT
DGLTPVEMQNLEPGTCYQVSGRCQVENGYPWGEWSSPLSFQTPFLDPEDVWVSGTVCETS
GKRAALLVWKDPRPCVQVTTYTVWFGAGDITTTQEEVPCCKSPVPAMMEWAVVSPGNSTSW
VPPTNLSLVCLAPESAPCDVGVSSADGSPGIKVTWKQGTTRKPLEYVVDWAQDGDSDKLN
WTRLPPGNLSTLLPGEFKGGVPYRITVTAVYSGGLAAAPSVWGFREELVPLAGPAVWRLP
DDPPGTPVVAWGEVPRHQLRGQATHYTFICIQSRGLSTVCRNVSSQTQTATLPNLHSGSFK
LWVTVSTVAGQGPPGPDLSLHLPDNRIWKALPWFLSLWGLLLMGCGLSLASTRCLQARC
LHWRHKLKPQWIWERPDPANSNSGQPYIKEVSLPQPPKDGPILEVEEVELQPVVESPKA
SAPIYSGYEKHFLPTPEELGLLV

FIG._4

ADULT													FETAL							
PBLs	Colon	Sm. Int.	Ovary	Testis	Prostate	Thymus	Spleen	Heart	Brain	Placenta	Lung	Liver	Sk. Muscle	Kidney	Pancreas	Kidney	Liver	Lung	Brain	Heart
																				

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FIG._6

APPROVED	O.G. FIG.	
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h-TCCR	1	MRGGRGAPFWLWLP	PKLALPLLWVLFQRTRP	QGSAGPLQCY	GVGPLGDL
m-TCCR	1	-----MNRRLRVARI	TPLLELLSLMSLLGTRPHGSP	GPLQCY	SVGPLGIL
h-TCCR	51	NCSWEPLGDLGAPSE	HLQSQKYSRNSKTQTVA	AAAGRSWVAIPREQ	LTMS
m-TCCR	46	NCSWEPLGDLLET	PPVLYHQSQKYHPNRVWE	VKPSKQSWVTIPREQ	FTMA
h-TCCR	101	DKLLVWGTAKAGQ	PLWPPVFNLEETQMKPNAP	RLGPDVD	FSEDDPLEATVH
m-TCCR	96	DKLLIWGTQKGR	PLWSSVSVNLEETQMKPD	TPIFSQVD	ISEEATLEATVQ
h-TCCR	151	WAPPTWPSHKVL	ICQFHYRRCCQEAAT	LLEPELKT	IPLTPVEIQDLELAT
m-TCCR	146	WAPPVWPPQKAL	TCQFRYKECQAEAT	RLLEPQLK	TDGLTPVEMQNLEPGT
h-TCCR	201	GKVVYGRCRMKEED	LWGEWSPILSFQTPPSA	PKDVWVSGNL	CGTTPGGEE
m-TCCR	195	CYQVSGRCQVENGY	P-WGEWSSPLSFQTP	FLDPEDEVVSG	TVCETSGKRA
h-TCCR	251	PIILLWKA	PGCVQVSQVWFFV	GGRELSPEGIT	CCCSLIPSGAEWARVSA
m-TCCR	245	ALLVWKD	PRPCVQVTTYVWF	GAGDITTTQEE	VPEVPCCKSPVPAWMEWAVVSP
h-TCCR	301	VNATSWEP	LTNLSLVCLDSASAP	RSVAVSS	IAGSTELLVTWQPGGEP
m-TCCR	295	GNSTSWVP	PTNLSLVCLAPESAP	CDVGVSSAD	GSPIKVTWKQGT
h-TCCR	351	HVVDWARD	GDPLEKLNWVRL	PPGNLSALL	PGNFTVGVYRITVTAVSASG
m-TCCR	345	YVVDWAQ	DGDSLDKLNWTR	RLPPGNLS	TLLPGEFKGGVYRITVTAVS
h-TCCR	401	LAASA	SSVWGFREELAP	LVGPTLWRLQDA	PPGTPIAWGEVPRHQLRGHLT
m-TCCR	395	LAAP	SVWGFREELVPL	AGPAVWRLP	DDPPGTPIAWGEVPRHQLRGQAT
h-TCCR	451	HYTLCAQ	SGTSPSVCMNV	SGNTQSVTL	PLPWGPCELWVTASTIAGQGPP
m-TCCR	445	HYTFCIQ	SRGLSTVCRNV	SSQTQTATLP	NLHSGSFKLWVTVSTVAGQGPP
h-TCCR	501	GPILLRLHLP	DNTRLRWKVL	PLPGILFLWGL	FLGCGGLSLATS---
m-TCCR	495	GPDL	SLHLPDNRIRWKA	LPWFLSLWGL	LLMCGGLSLASTRCLQARCLHWR
h-TCCR	547	HKVLP	PRWVWEKVPDP	PANSSSGQPHMEQ	VPEAQLGDLFILEVEEEMEPPPV
m-TCCR	545	HKLLPQW	IWERVDPDPANS	NSGQPYIKEVSL	PQPKDGPFILEVEEVEQLQPV
h-TCCR	597	MES	SQPAQATAPLD	SGYEKHF	LPTPEELGLGPPRPQVLA
m-TCCR	595	VES	-----PKASAPI	YSGYEKHF	LPTPEELGLV

Box 1

FIG.-5





APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

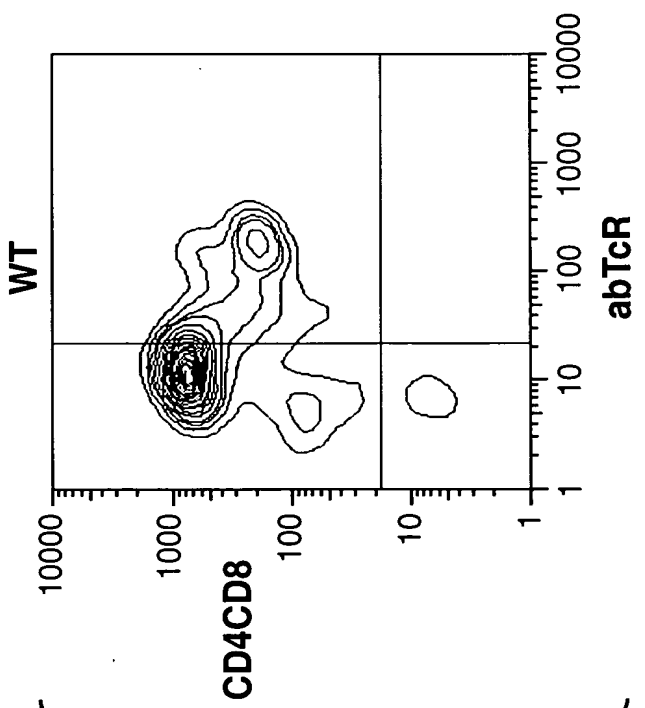
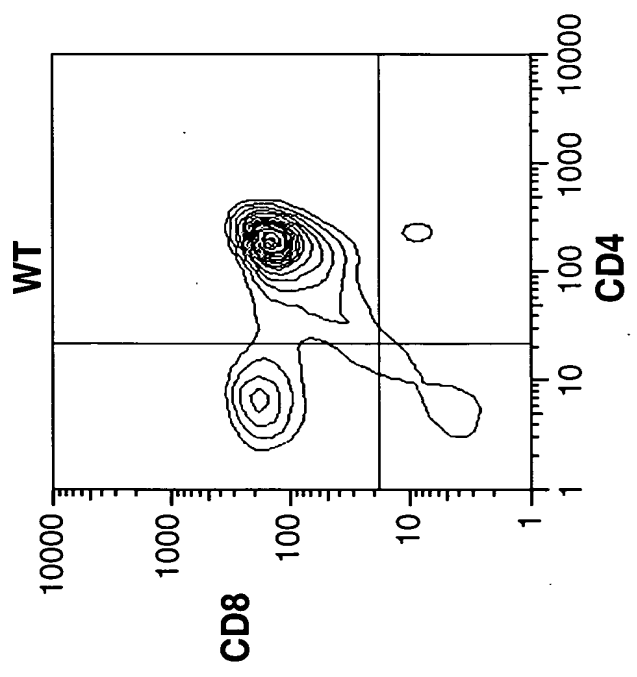
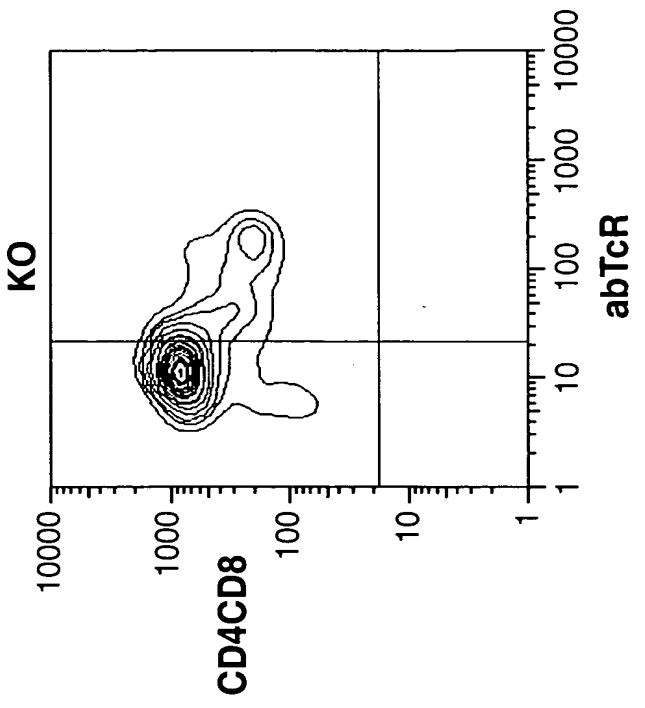
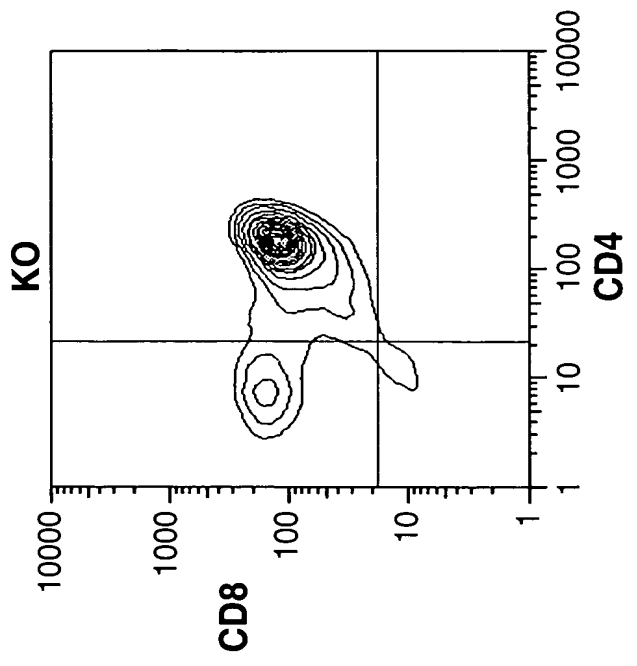


FIG.-7A

FIG.-7B

APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

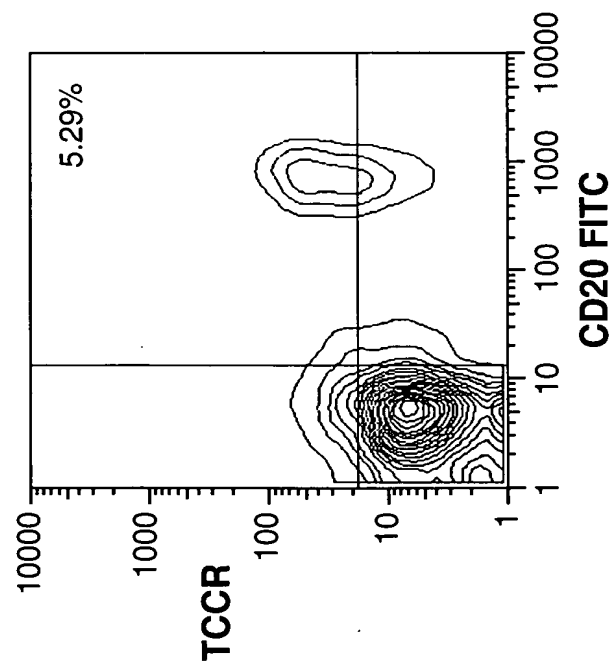
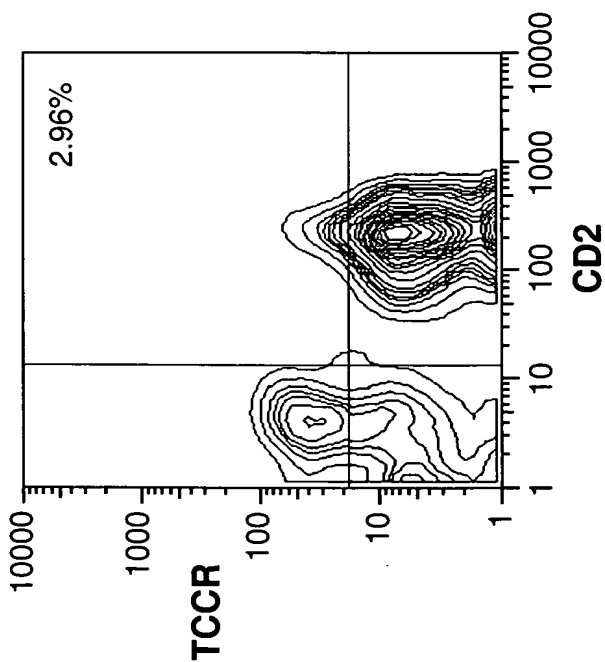
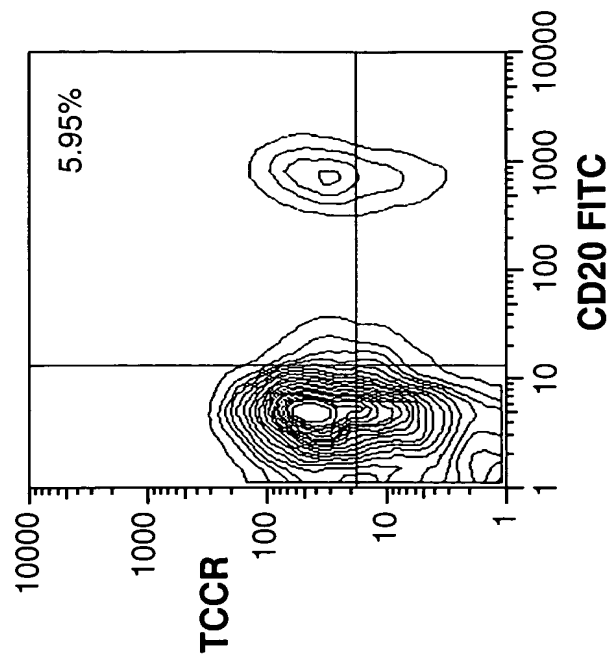
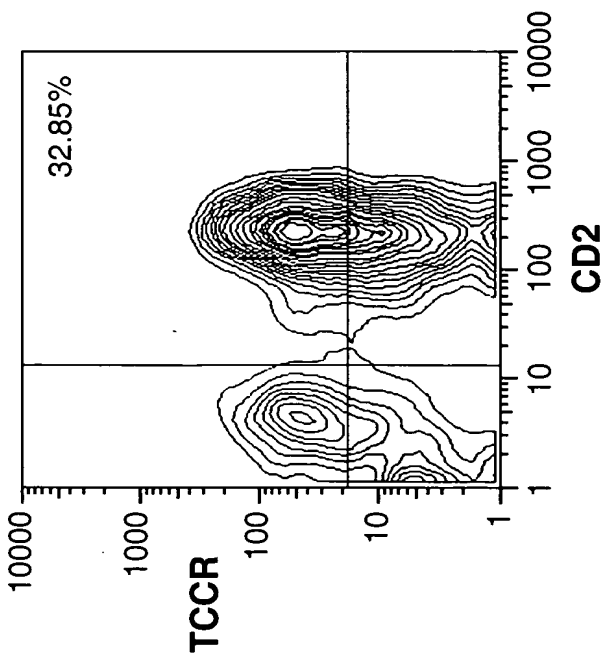


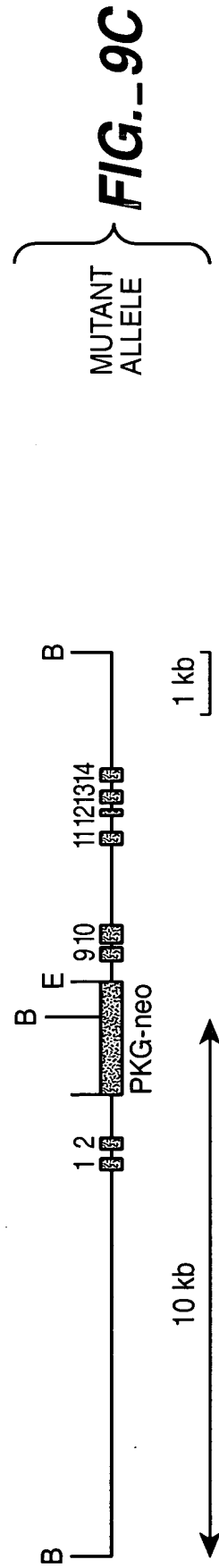
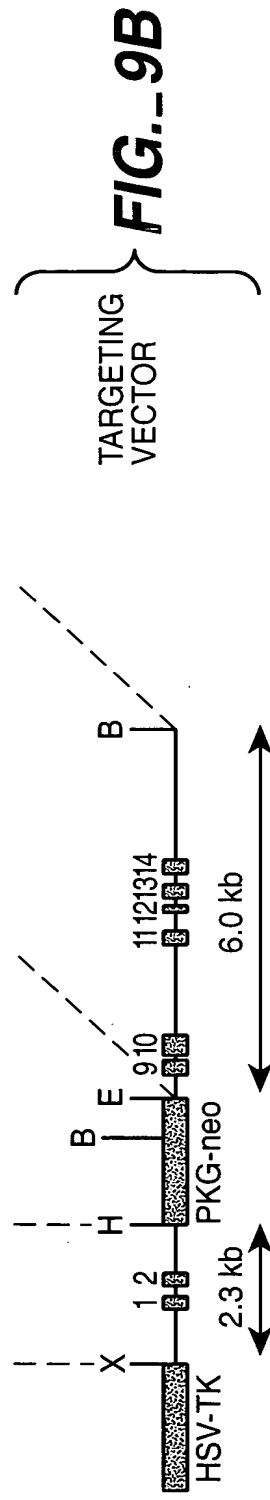
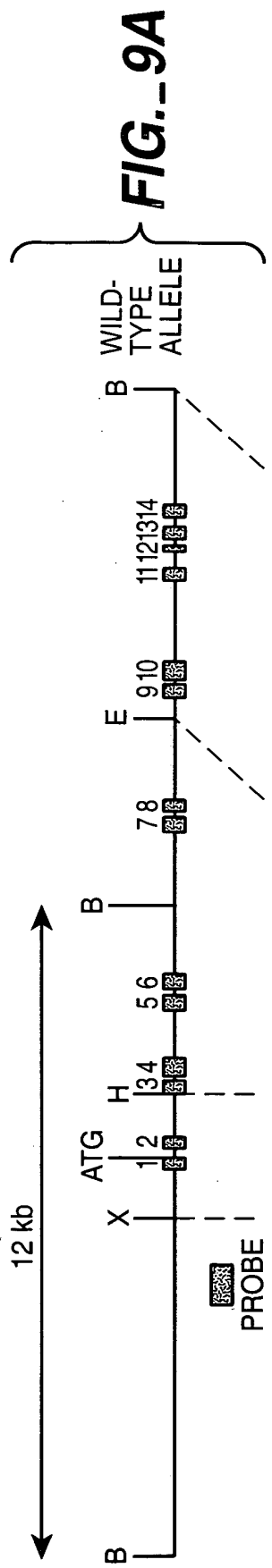
FIG._8A

FIG._8B



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O.G. FIG.		
APPROVED	CLASS	SUBCLASS
BY		
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BY	CLASS	SUBCLASS
DRAFTSMAN		

FIG._10A

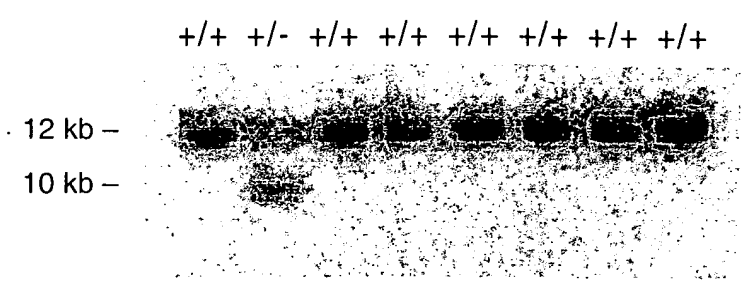


FIG._10B

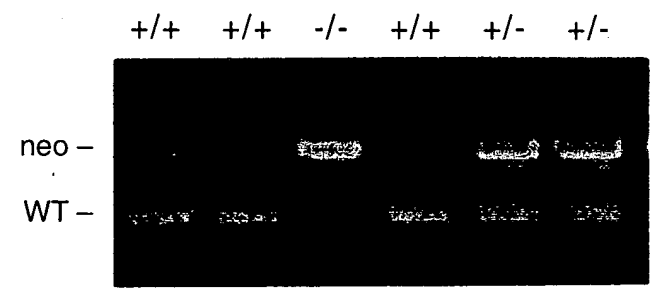
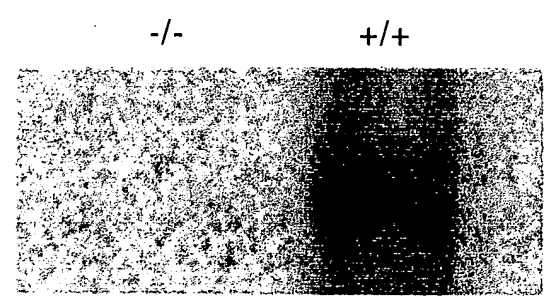


FIG._10C



APPROVED BY	O.G. FIG.		SUBCLASS
	CLASS		
DRAFTSMAN			

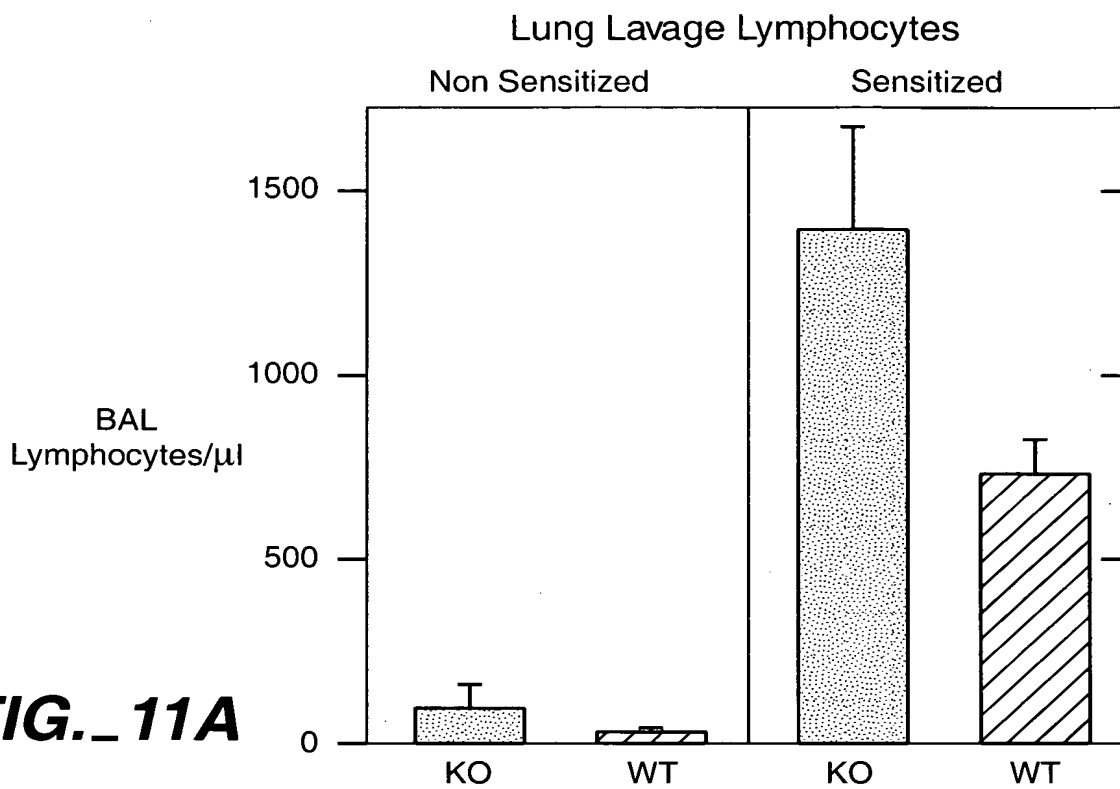


FIG._11A

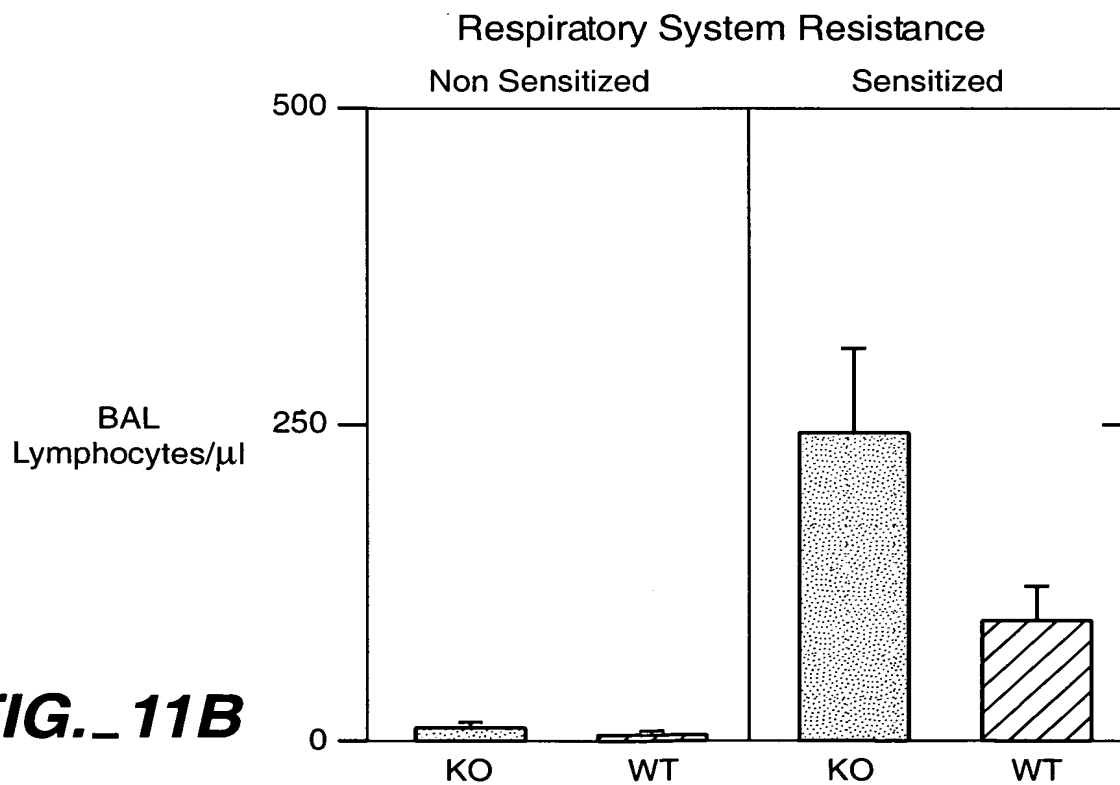


FIG._11B

APPROVED	O.G. FIG.	SUBCLASS
BY	CLASS	
DRAFTSMAN		

FIG._12A

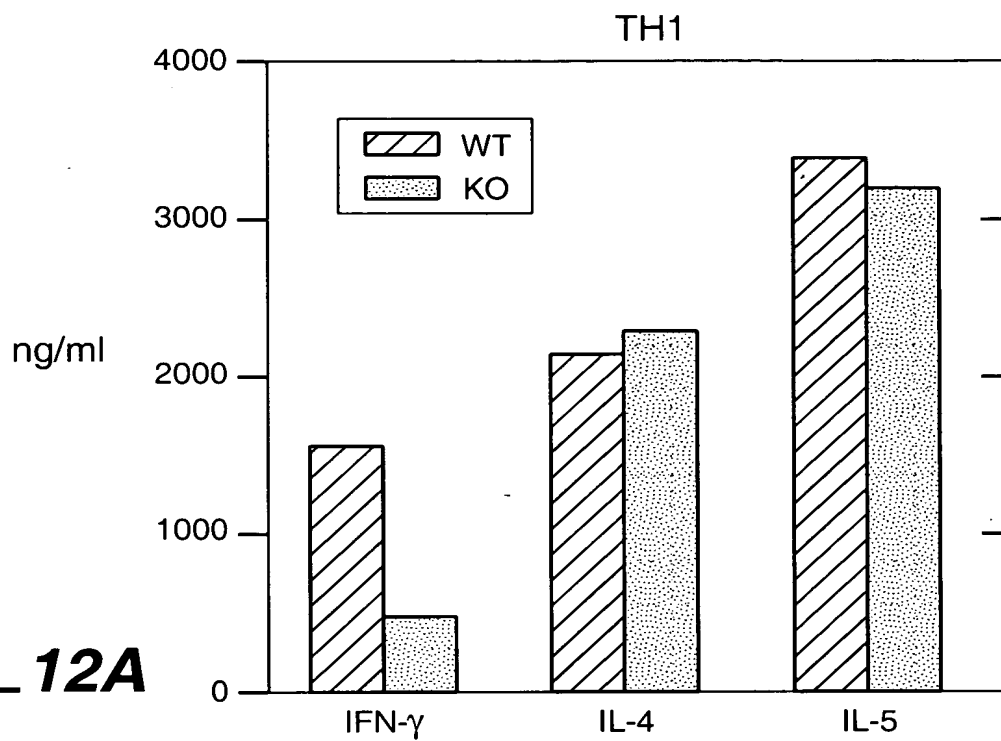
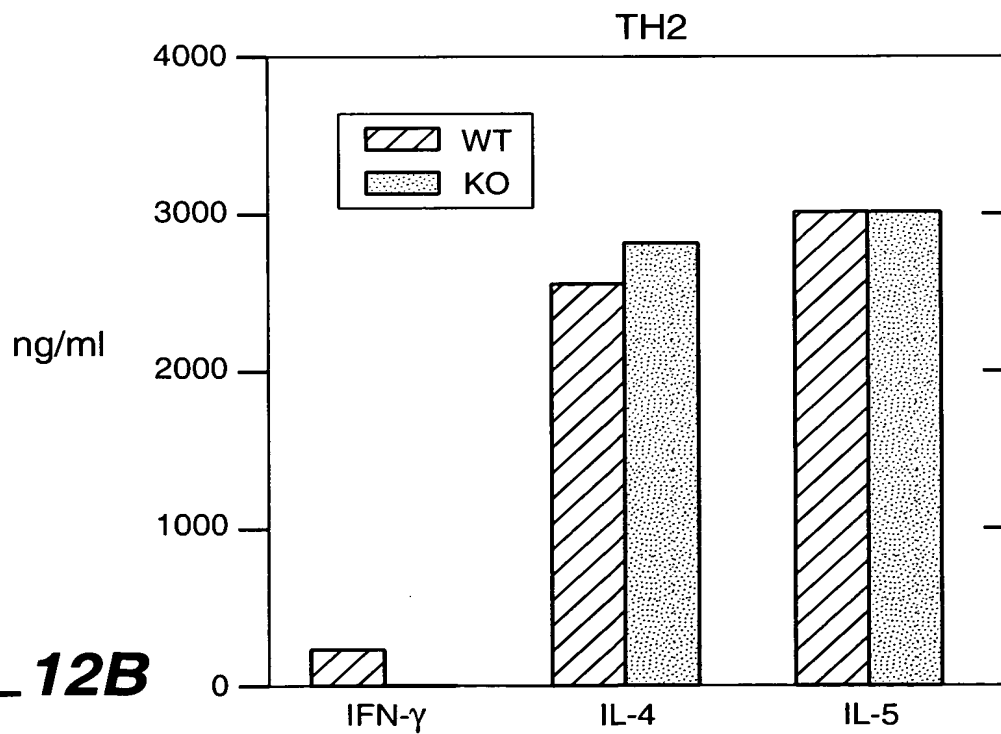


FIG._12B



APPROVED BY DRAFTSMAN	O.G. FIG.	
	CLASS	SUBCLASS

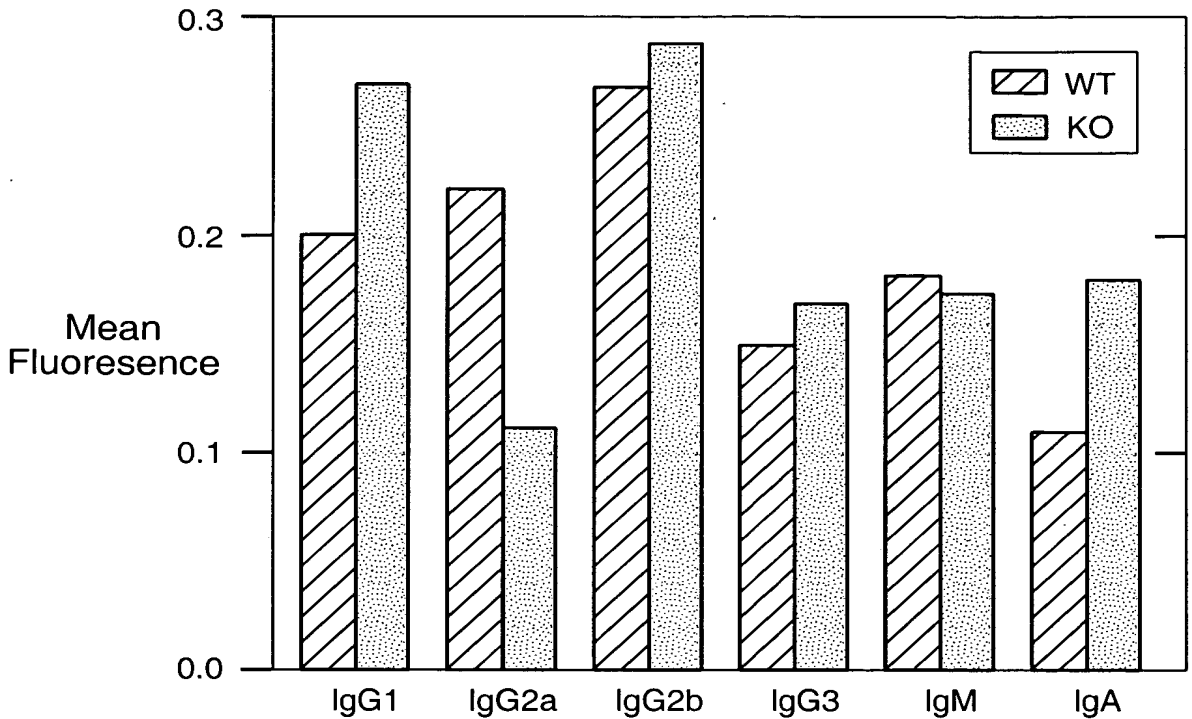


FIG._13

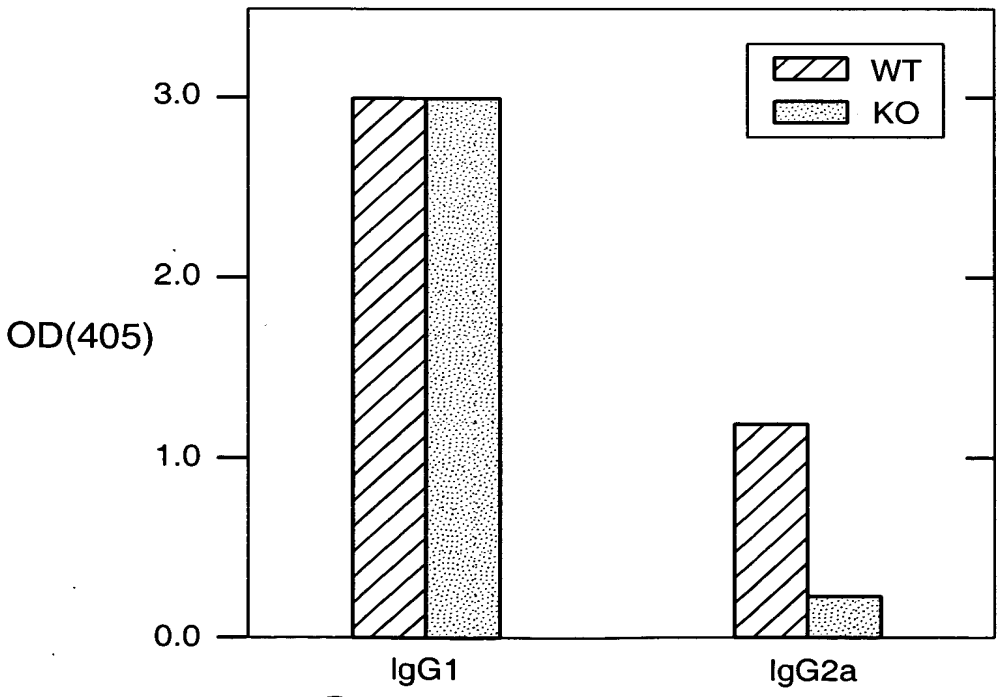


FIG._14

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APPROVED BY DRAFTSMAN	O.G. FIG.	SUBCLASS
	CLASS	

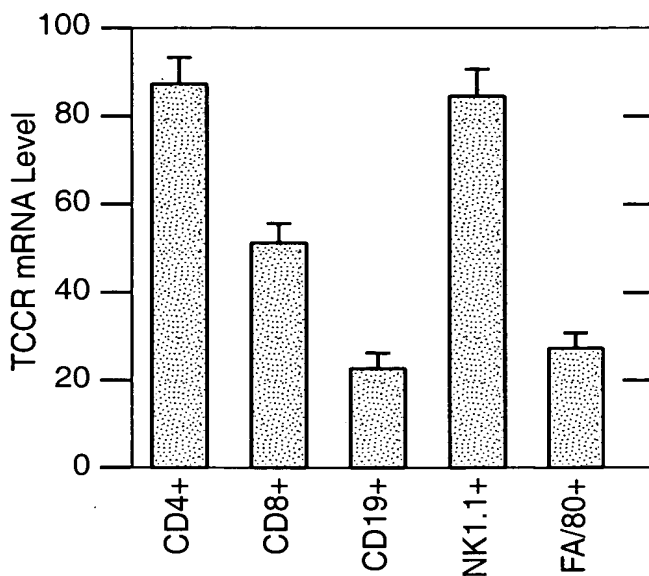


FIG._15A

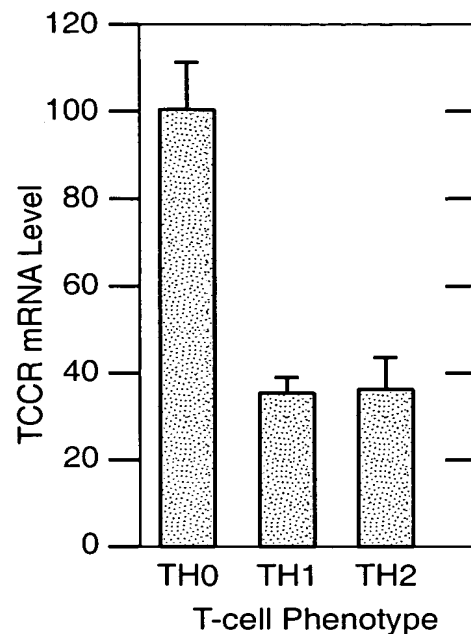


FIG._15B

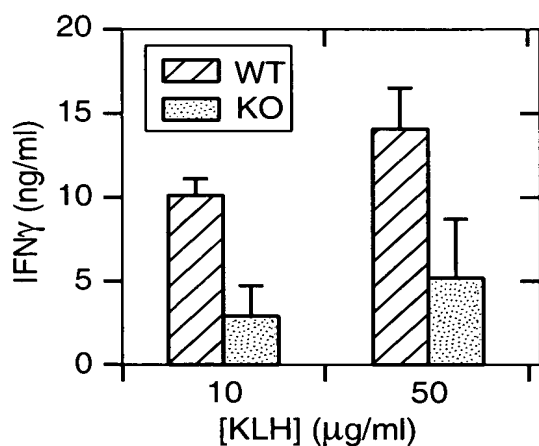


FIG._16A

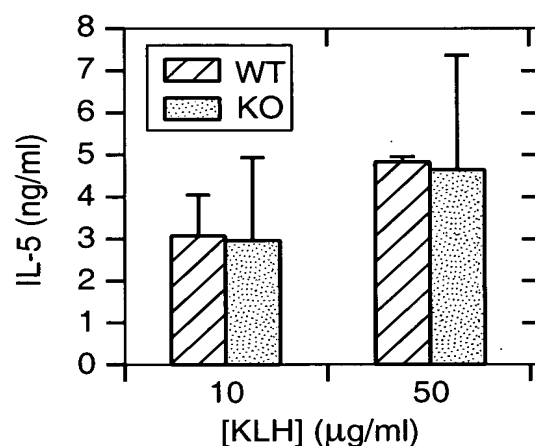


FIG._16C

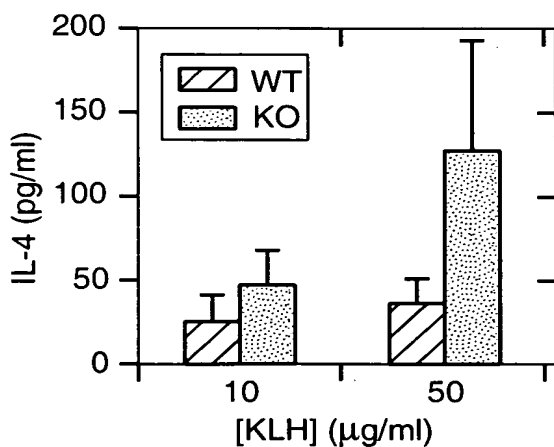


FIG._16B

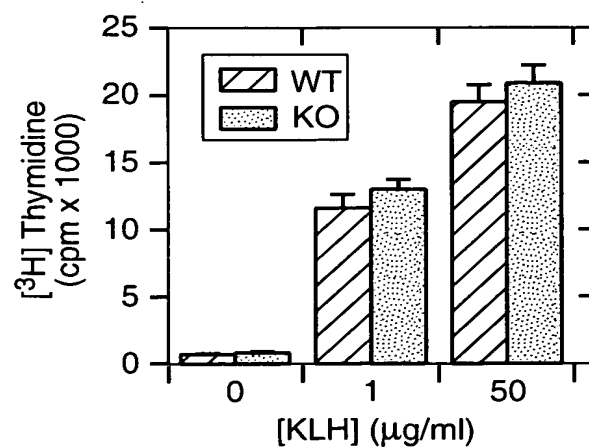


FIG._16D

APPROVED BY DRAFTSMAN	O.G. FIG.	SUBCLASS
	CLASS	

FIG._17A

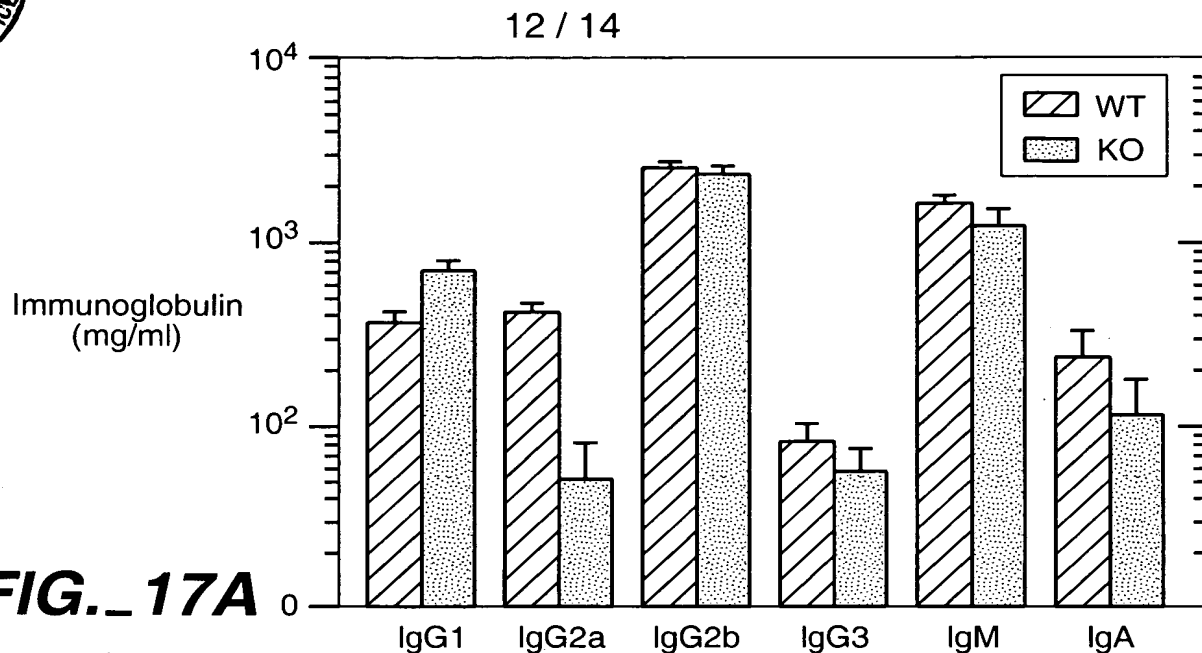


FIG._17B

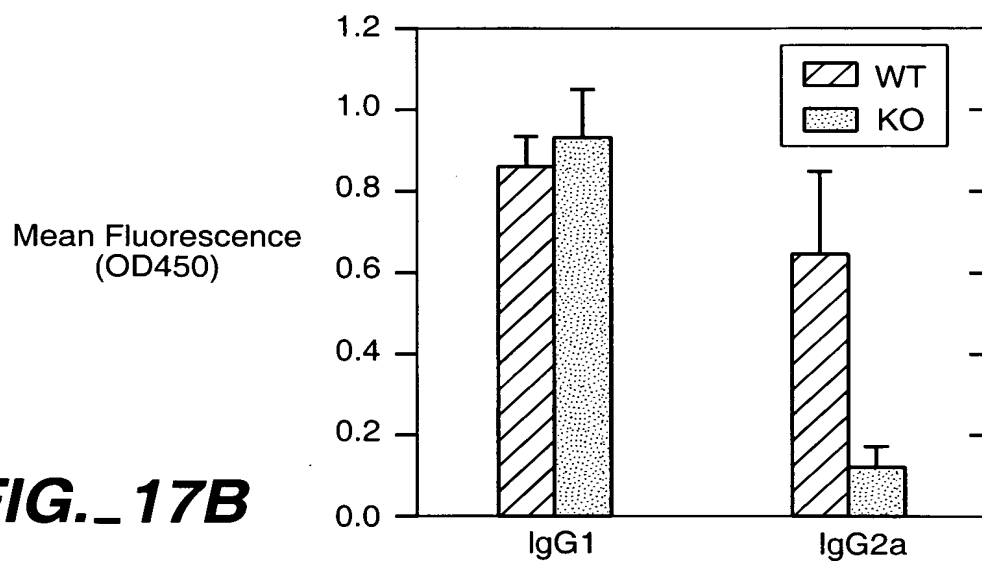
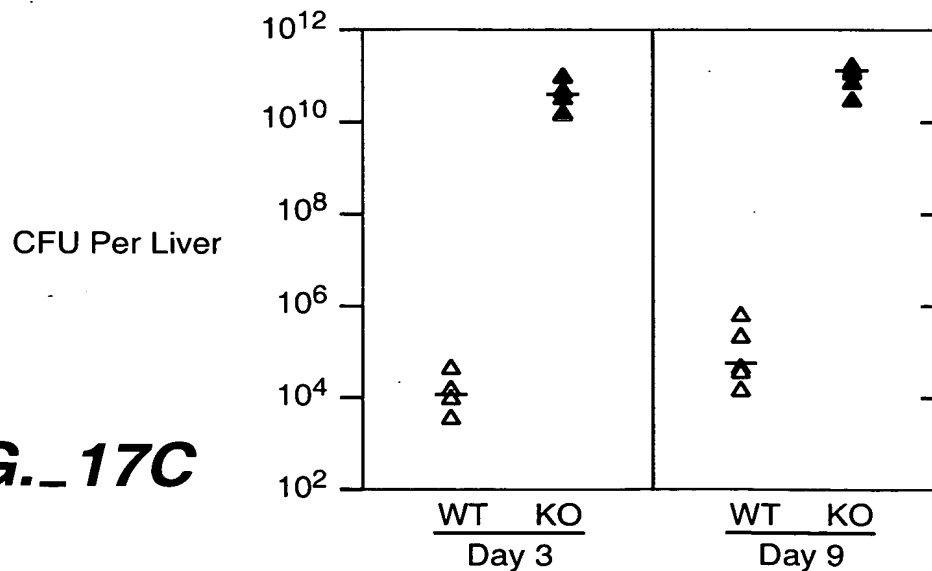


FIG._17C



APPROVED BY DRAFTSMAN	O.G. FIG.	SUBCLASS
	CLASS	

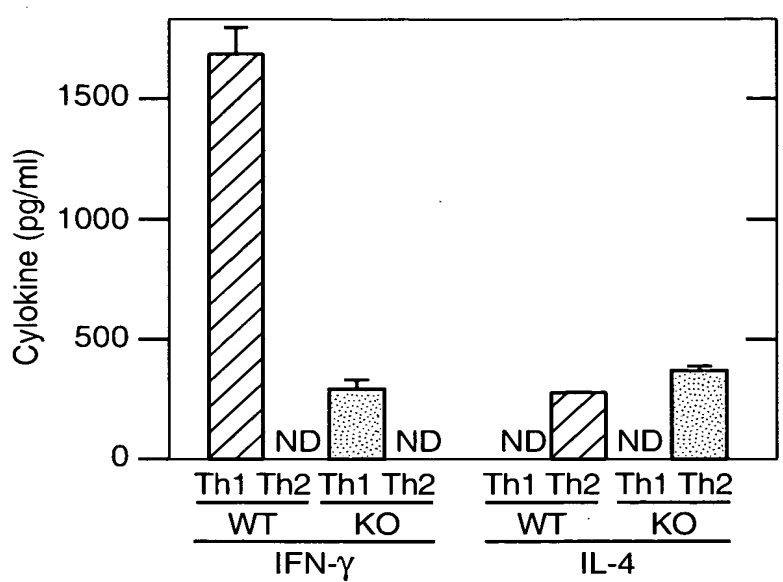


FIG._18A

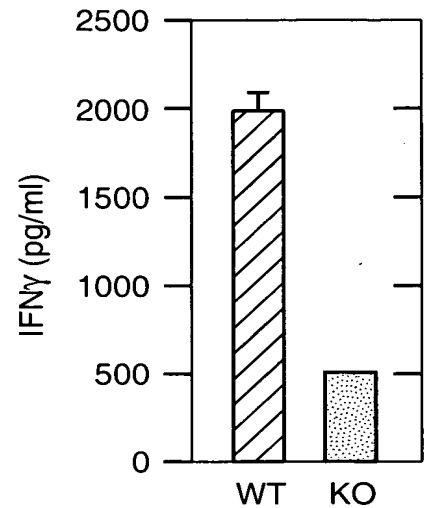


FIG._18B

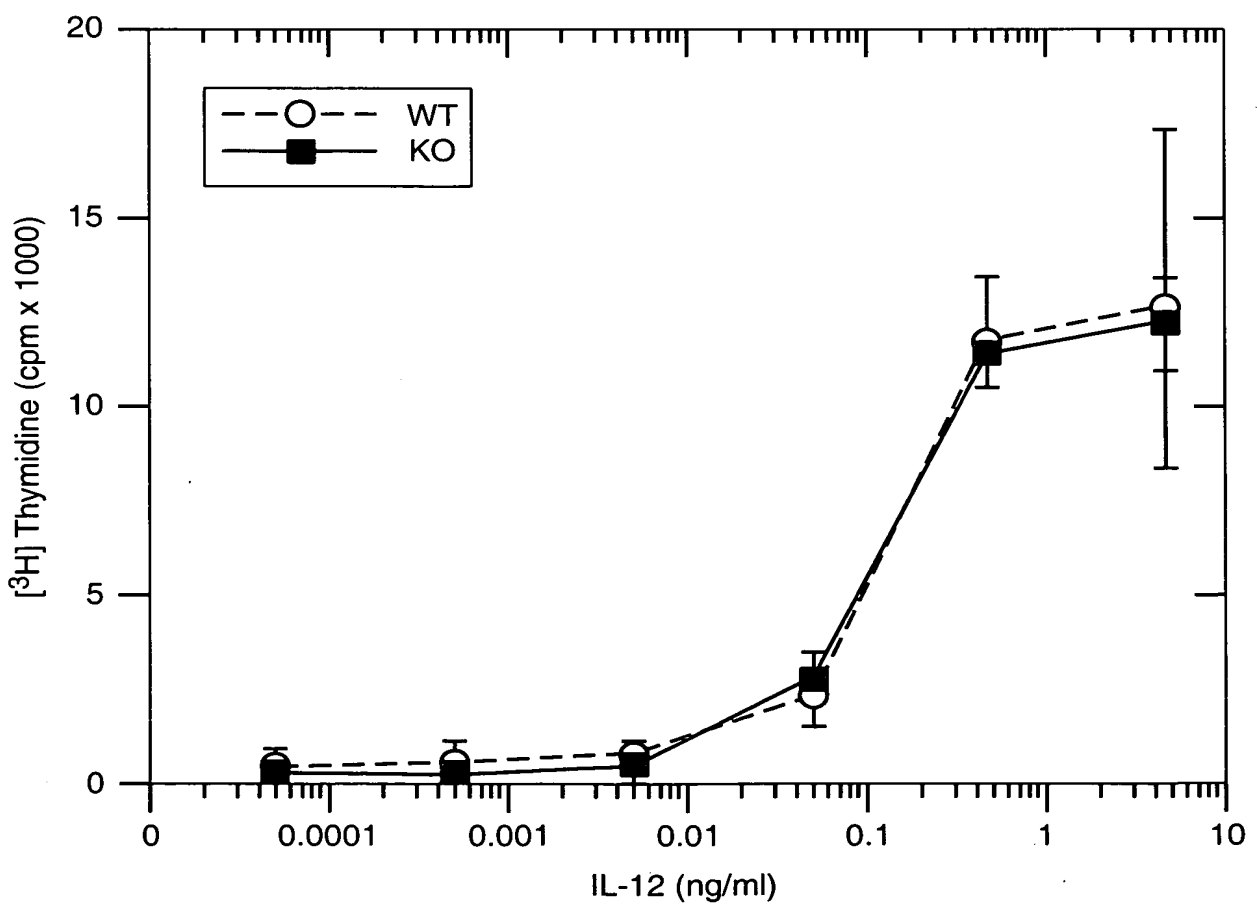


FIG._18C

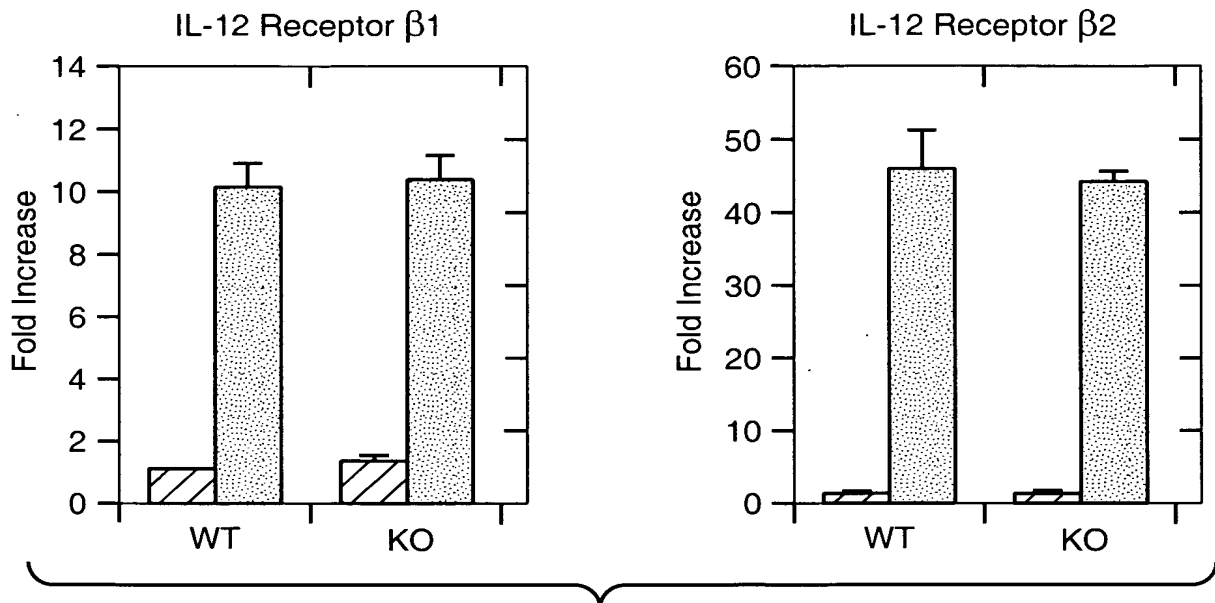


FIG._18D

Primer / Probe	Sequence	SEQ ID NO:
mTCCR, sense, Taqman	TGGTCTCTCCTGGCAACAGC	5
mTCCR, as, Taqman	AGCCAAGCACACCAGAGACA	6
mTCCR, Taqman probe	CAGCTGGGTGCCTCCCACCAA	7
mRPL19, sense, Taqman	ATCCGCAAGCCTGTGACTGT	8
mRPL19, as, Taqman	TCGGGCCAGGGTGTTTTT	9
mRPL19, Taqman probe	TTCCCGGGCTCGTTGCCG	10
mIL12Rb1, sense, Taqman	TCGCGTCTCTGGGAAGCT	11
mIL12Rb1, as, Taqman	TTTAAGCCAATGTATCCGAGACTG	12
mIL12Rb1, Taqman probe	CGCCAGCGTCCTCCTCGTGG	13
mIL12Rb2, sense, Taqman	CAAGCATTTGCATCGCTATCA	14
mIL12Rb2, as, Taqman	AATGCCTTTTGCCGGAAGT	15
mIL12Rb2, Taqman probe	ACGAATTGAGAACGTGCCCACCGT	16

FIG._19